

FIG. I

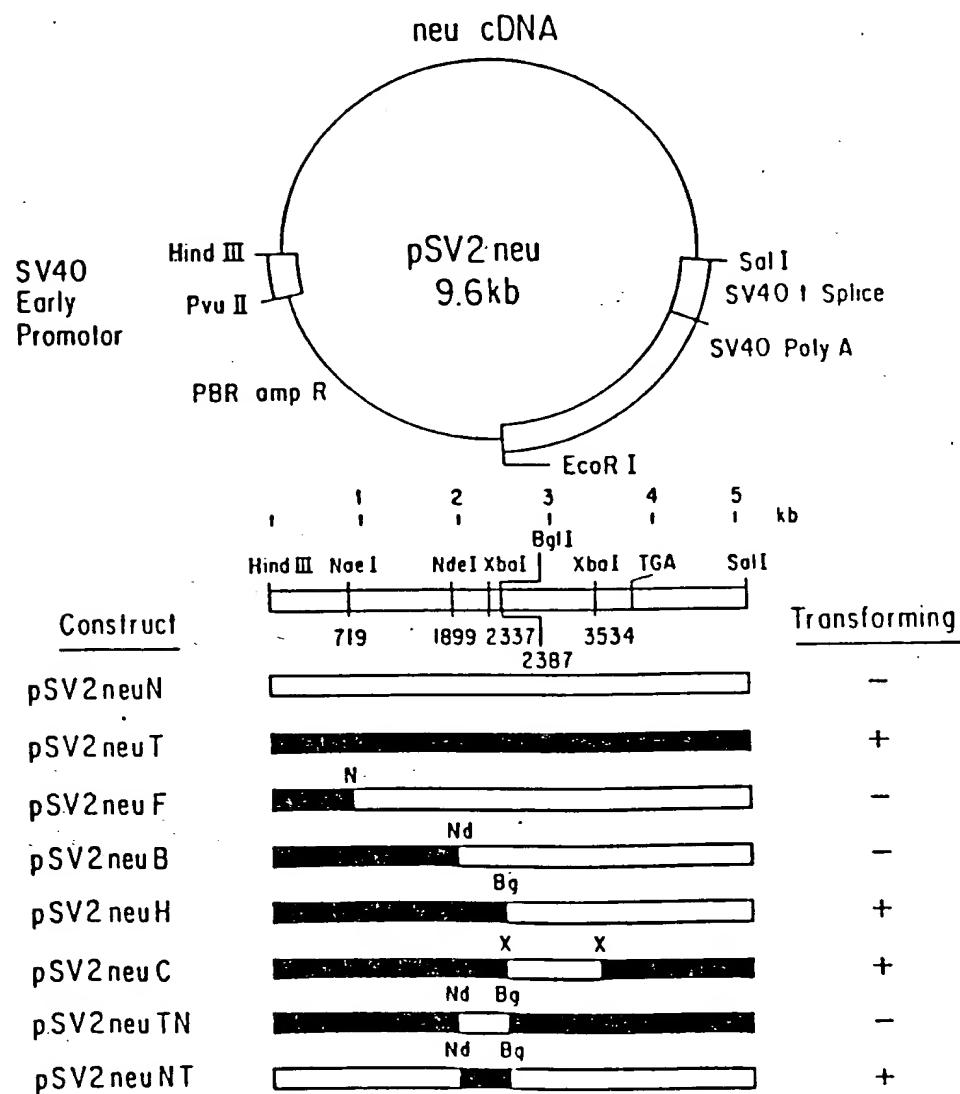
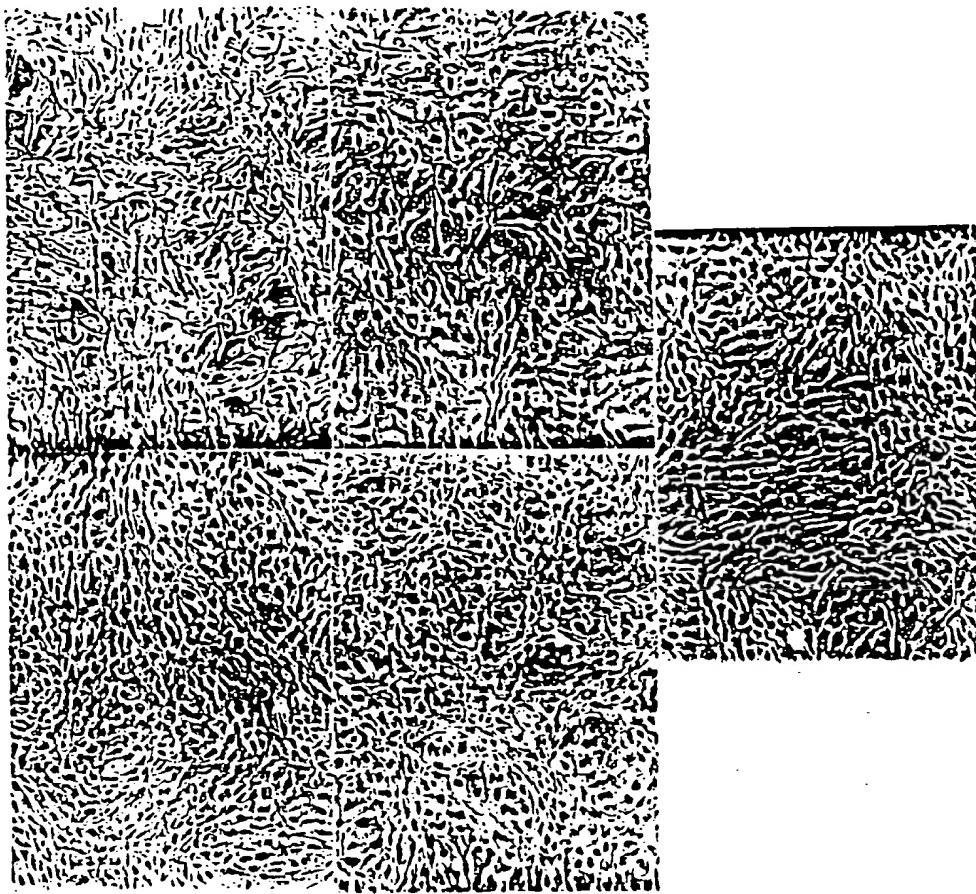
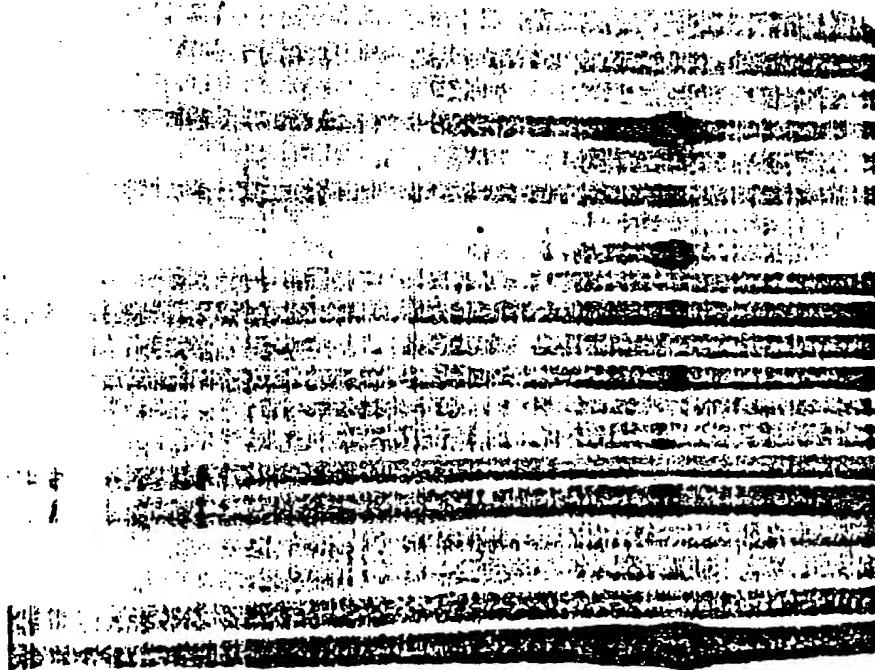


FIG.2A



p 185 —

FIG.2B



a b c d e f g h i j
- + - + - + - + - + - + - +

FIG.3

| | | | | |
|---|---|---------|--|--------|
| normal | | val | | |
| | | GTG | | |
| glu gln arg ala ser pro val thr phe ile ile ala thr val | : | gly val | | aa 666 |
| GAG CAG AGA GCC AGC CCG GTG ACA TTC ATC ATT GCA ACT GTA | : | GGC GTC | | |
| | | CAG | | |
| transforming | | glu | | |
| | | | | |
| leu leu phe leu ile leu val val val val gly ile leu ile lys arg arg | | | | aa 683 |
| CTG CTG TTC CTG ATC TTA GTG GTG GTC GTT GGA ATC CTA ATC AAA CGA AGG | | | | |

FIG.4

- A) ACGCCCAC TACAGTTGCAAT nucleotides 1999-2018, wild-type sequence
- *
B) ACGCCCTCTACAGTTGCAAT nucleotides 999-2018, T₂₀₁₂ to A
- C) CCGTCCTCAGCTGTGACC nucleotides 996-1013, control probe
- *
D) ACGCCCCCTACAGTTGCAAT nucleotides 1999-2018, T₂₀₁₂ to G

FIG.5

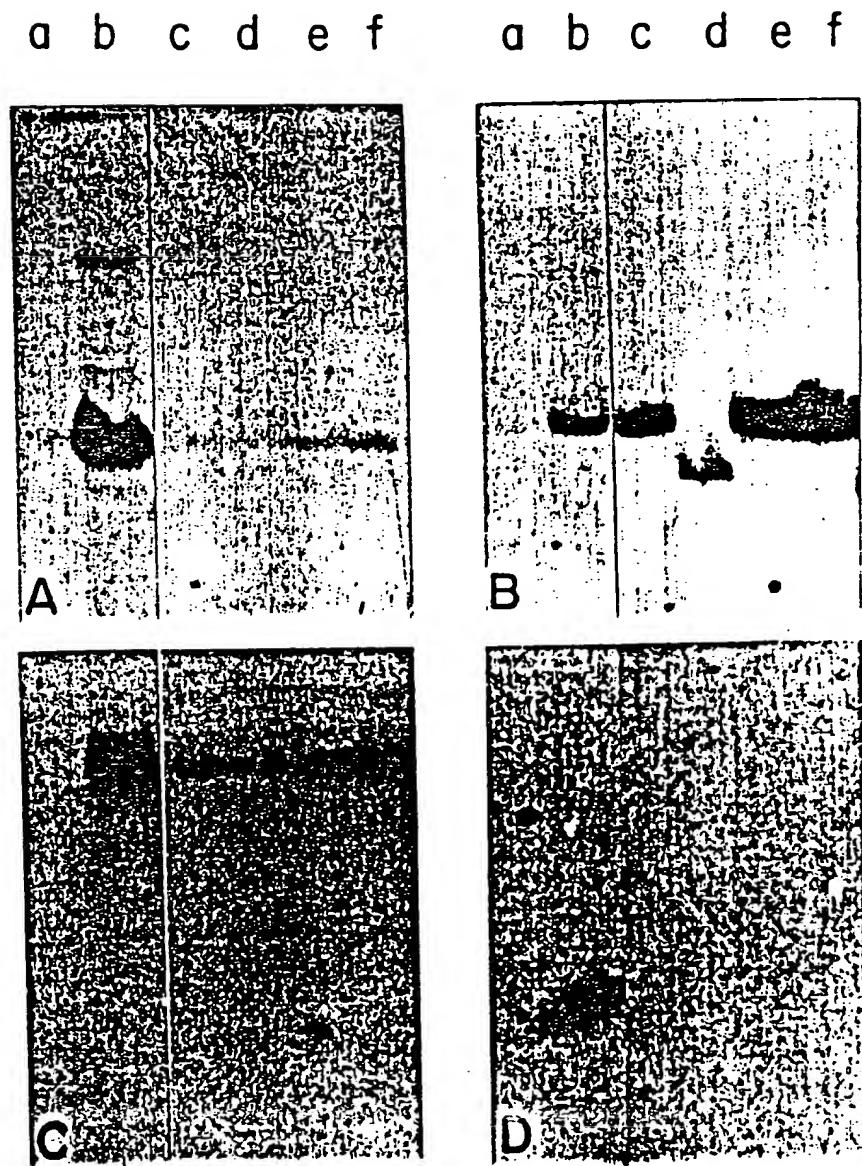


FIG. 6

a b c d e f g h i j

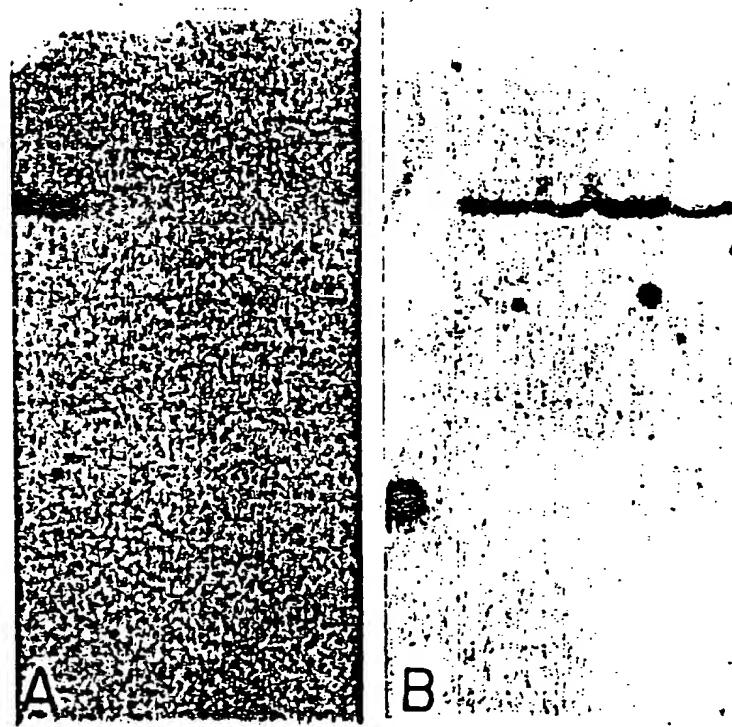
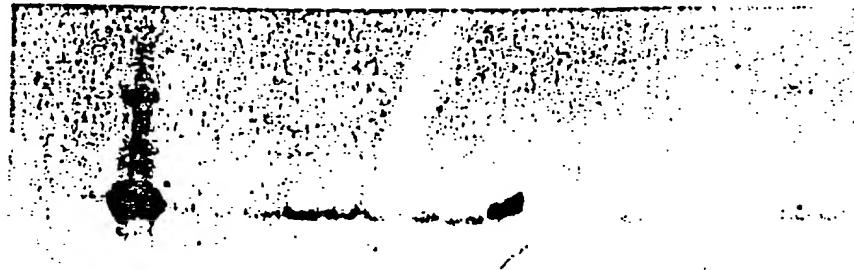
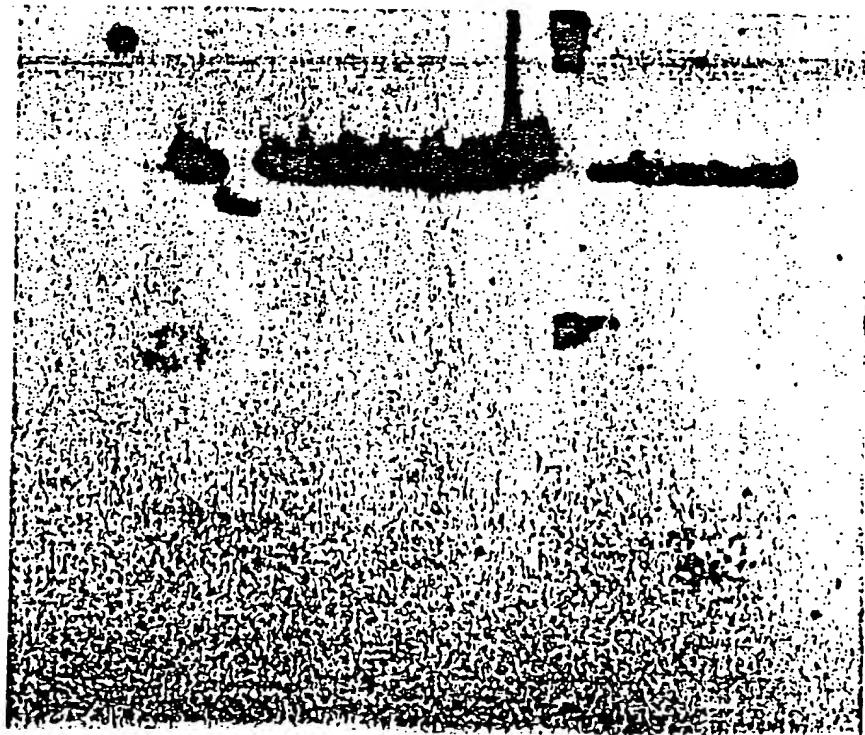


FIG.7

N G B 1 2 3 4 5 6 7 8 9 10 11 12 13 14 c1 c2



N G B 1 2 3 4 5 6 7 8 9 10 11 12 13 14 c1 c2



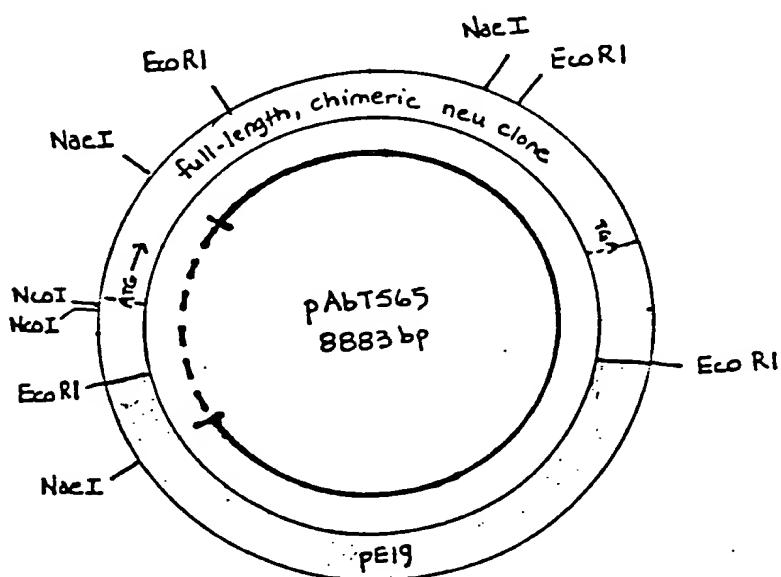
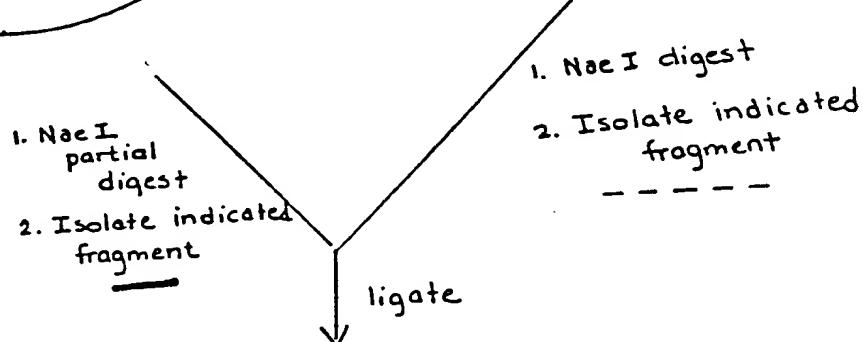
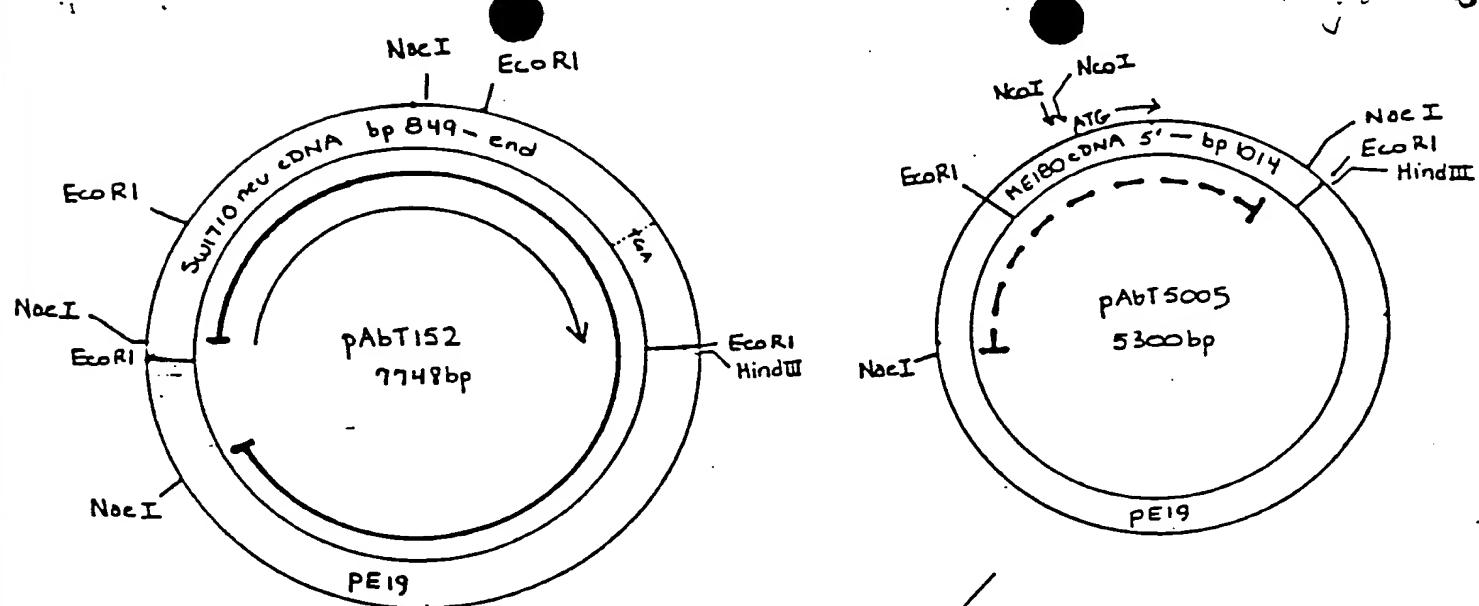


Fig 11. Construction of full length-neu cDNA clone from ME180 and SW1710 neu cDNAs

Construction of pMax neu

Figure 9

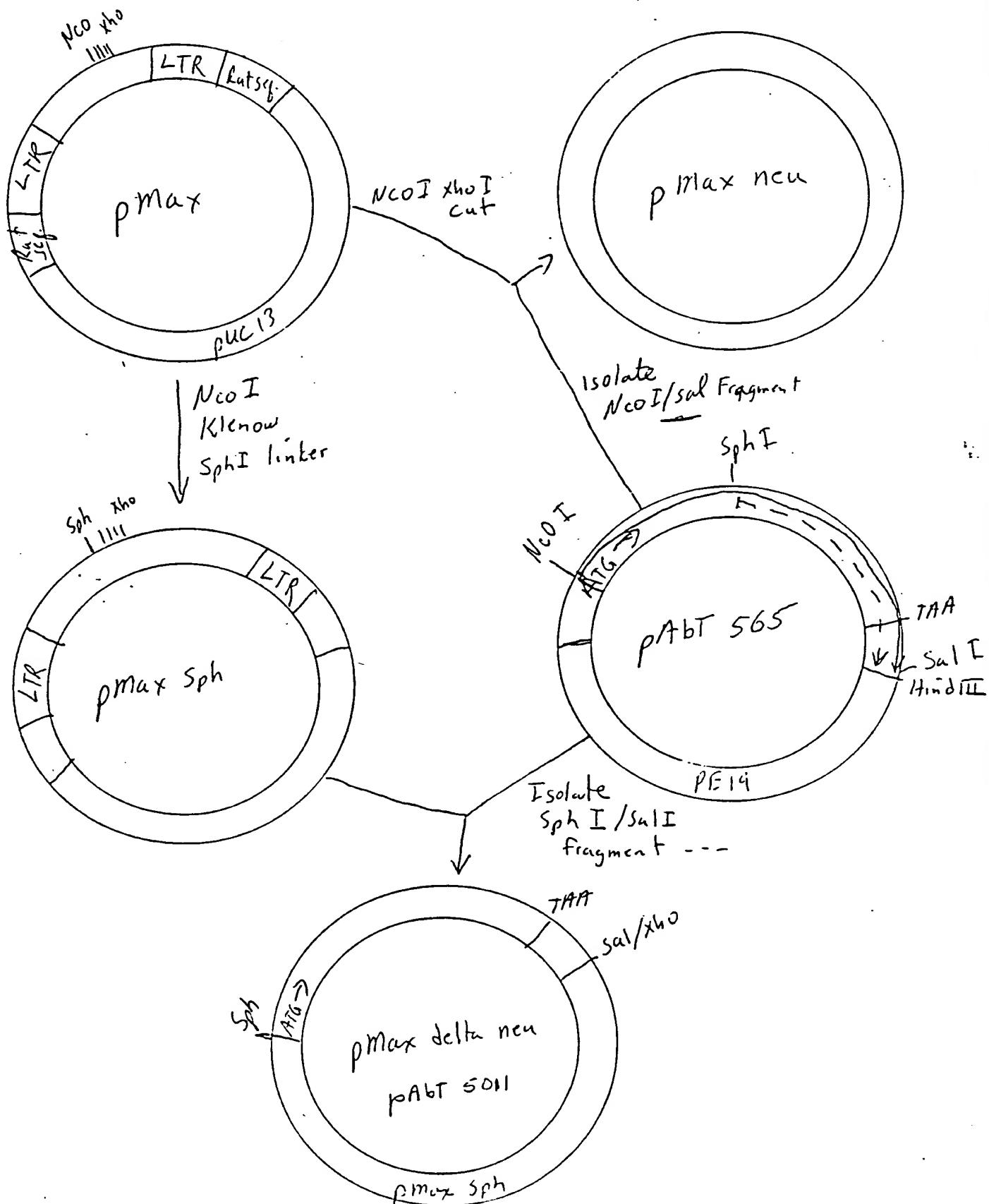


Figure 114
Neu Oncogene Probe Design:

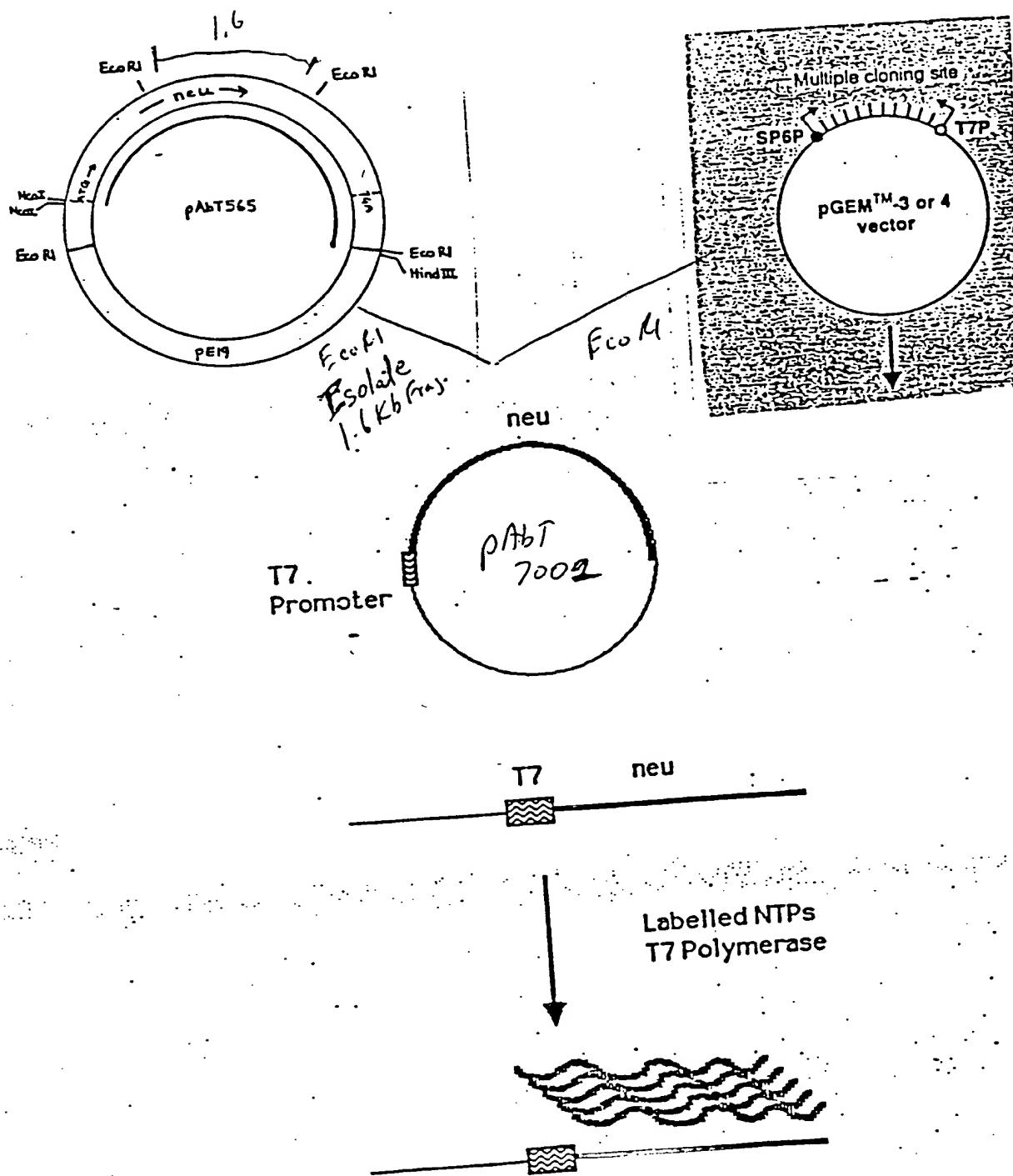
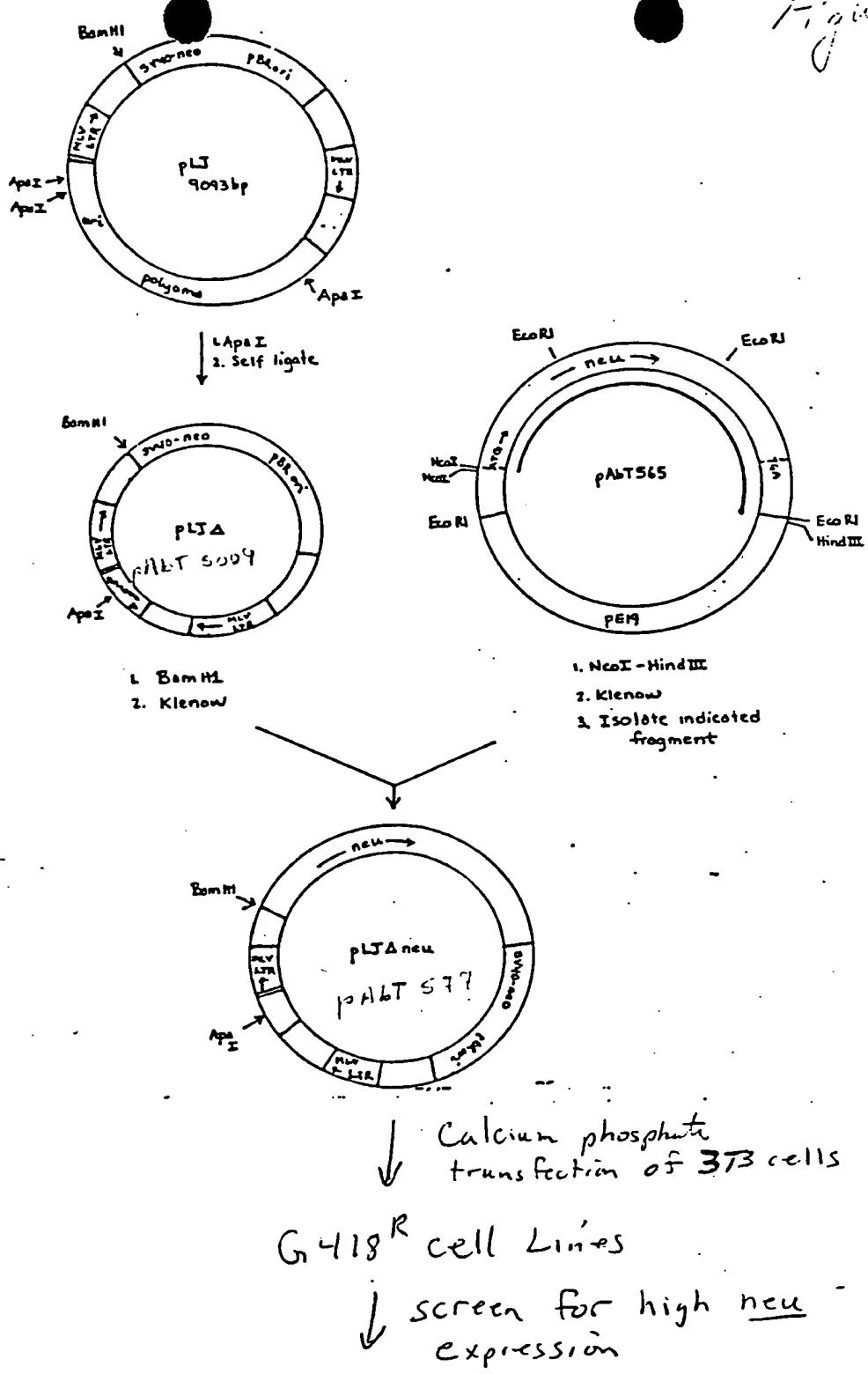


Figure 11



18-3-7